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Thr Thr Ala Leu Arg Ala Leu Asn Gly Leu Gly Cys Ala Gly Val Pro

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Gly Glu Thr Ala Gly Gly Ala Val Gly Pro Gly Pro Leu Gly Thr Arg

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Gly Phe Leu Ser Gly Ser Lys Phe Gln Ala Pro Gly Ser Trp Lys Asp

65                    70                    75                    80

Cys Phe Gly Ala Pro Pro Ala Pro Asp Val Leu Arg Ala Asp Arg Ser

                  85                    90                    95

Val Gly Glu Gly Cys Pro Gln Lys Leu Val Thr Ala Asn Leu Leu Arg

                  100                    105                    110

Phe Leu Leu Leu Val Leu Ile Pro Cys Ile Cys Ala Leu Ile Val Leu

                  115                    120                    125

Leu Ala Ile Leu Leu Ser Phe Val Gly Thr Leu Lys Arg Val Tyr Phe

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| Pro Ser Leu Pro Pro Ser Gln Ser Thr Pro Ala Trp Thr Pro Arg Ala |  |     |  |     |
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| Pro Ser Pro Glu Asp Gln Ser His Arg Asn Thr Ser Thr Cys Met Asn |  |     |  |     |
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| Phe Gly Cys Ser Leu Ala Phe Pro Glu Cys Val Val Asp Gly Asp Asp |  |     |  |     |
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Arg Lys Ser Cys Phe Ser Leu Gln Gln Glu His Gly Lys Gln Ser Leu

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Cys Gly Gly Gly Glu Ser Phe Leu Cys Thr Ser Gly Leu Cys Val Pro

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Lys Lys Leu Gln Cys Asn Gly Tyr Asn Asp Cys Asp Asp Trp Ser Asp

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360

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Glu Ala His Cys Asn Cys Ser Lys Asp Leu Phe His Cys Gly Thr Gly

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375

380

Lys Cys Leu His Tyr Ser Leu Leu Cys Asp Gly Tyr Asp Asp Cys Gly

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390

395

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Asp Pro Ser Asp Glu Gln Asn Cys Asp Cys Asn Leu Thr Lys Glu His

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Arg Cys Gly Asp Gly Arg Cys Ile Ala Ala Glu Trp Val Cys Asp Gly

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Phe Gly Ser Ser Cys Val Glu Ser Cys Ala Gly Ser Ser Leu Cys Asp

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895

Cys Val Leu Ile Ala Lys Lys Trp Val Leu Thr Val Ala His Cys Phe

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Glu Gly Arg Glu Asp Ala Asp Val Trp Lys Val Val Phe Gly Ile Asn

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Asn Leu Asp His Pro Ser Gly Phe Met Gln Thr Arg Phe Val Lys Thr

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Gly Leu Thr Ser Trp Gly Ser Val Cys Phe Ser Lys Val Leu Gly

1070

1075

1080

Pro Gly Val Tyr Ser Asn Val Ser Tyr Phe Val Gly Trp Ile Glu

1085

1090

1095

Arg Gln Ile Tyr Ile Gln Thr Phe Leu Gln Lys Lys Ser Gln Gly  
 1100 1105 1110

<210> 4

<211> 1042

<212> PRT

<213> Homo sapiens

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Met Lys Gln Ser Pro Ala Leu Ala Pro Glu Glu Arg Tyr Arg Arg Ala  
 1 5 10 15

Gly Ser Pro Lys Pro Val Leu Arg Ala Asp Asp Asn Asn Met Gly Asn  
 20 25 30

Gly Cys Ser Gln Lys Leu Ala Thr Ala Asn Leu Leu Arg Phe Leu Leu  
 35 40 45

Leu Val Leu Ile Pro Cys Ile Cys Ala Leu Val Leu Leu Val Ile  
 50 55 60

Leu Leu Ser Tyr Val Gly Thr Leu Gln Lys Val Tyr Phe Lys Ser Asn  
 65 70 75 80

Gly Ser Glu Pro Leu Val Thr Asp Gly Glu Ile Gln Gly Ser Asp Val

85

90

95

Ile Leu Thr Asn Thr Ile Tyr Asn Gln Ser Thr Val Val Ser Thr Ala  
 100 105 110

His Pro Asp Gln His Val Pro Ala Trp Thr Thr Asp Ala Ser Leu Pro  
 115 120 125

Gly Asp Gln Ser His Arg Asn Thr Ser Ala Cys Met Asn Ile Thr His  
 130 135 140

Ser Gln Cys Gln Met Leu Pro Tyr His Ala Thr Leu Thr Pro Leu Leu  
 145 150 155 160

Ser Val Val Arg Asn Met Glu Met Glu Lys Phe Leu Lys Phe Phe Thr  
 165 170 175

Tyr Leu His Arg Leu Ser Cys Tyr Gln His Ile Met Leu Phe Gly Cys  
 180 185 190

Thr Leu Ala Phe Pro Glu Cys Ile Ile Asp Gly Asp Asp Ser His Gly  
 195 200 205

Leu Leu Pro Cys Arg Ser Phe Cys Glu Ala Ala Lys Glu Gly Cys Glu  
 210 215 220





Val Asp Lys Ser Asp Glu Val Asn Cys Ser Cys His Ser Gln Gly Leu

370

375

380

Val Glu Cys Arg Asn Gly Gln Cys Ile Pro Ser Thr Phe Gln Cys Asp

385

390

395

400

Gly Asp Glu Asp Cys Lys Asp Gly Ser Asp Glu Glu Asn Cys Ser Val

405

410

415

Ile Gln Thr Ser Cys Gln Glu Gly Asp Gln Arg Cys Leu Tyr Asn Pro

420

425

430

Cys Leu Asp Ser Cys Gly Gly Ser Ser Leu Cys Asp Pro Asn Asn Ser

435

440

445

Leu Asn Asn Cys Ser Gln Cys Glu Pro Ile Thr Leu Glu Leu Cys Met

450

455

460

Asn Leu Pro Tyr Asn Ser Thr Ser Tyr Pro Asn Tyr Phe Gly His Arg

465

470

475

480

Thr Gln Lys Glu Ala Ser Ile Ser Trp Glu Ser Ser Leu Phe Pro Ala

485

490

495

Leu Val Gln Thr Asn Cys Tyr Lys Tyr Leu Met Phe Phe Ser Cys Thr

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500

505

510

Ile Leu Val Pro Lys Cys Asp Val Asn Thr Gly Glu Arg Ile Pro Pro

515

520

525

Cys Arg Ala Leu Cys Glu His Ser Lys Glu Arg Cys Glu Ser Val Leu

530

535

540

Gly Ile Val Gly Leu Gln Trp Pro Glu Asp Thr Asp Cys Ser Gln Phe

545

550

555

560

Pro Glu Glu Asn Ser Asp Asn Gln Thr Cys Leu Met Pro Asp Glu Tyr

565

570

575

Val Glu Glu Cys Ser Pro Ser His Phe Lys Cys Arg Ser Gly Gln Cys

580

585

590

Val Leu Ala Ser Arg Arg Cys Asp Gly Gln Ala Asp Cys Asp Asp Asp

595

600

605

Ser Asp Glu Glu Asn Cys Gly Cys Lys Glu Arg Asp Leu Trp Glu Cys

610

615

620

Pro Ser Asn Lys Gln Cys Leu Lys His Thr Val Ile Cys Asp Gly Phe

625

630

635

640

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Pro Asp Cys Pro Asp Tyr Met Asp Glu Lys Asn Cys Ser Phe Cys Gln

645

650

655

Asp Asp Glu Leu Glu Cys Ala Asn His Ala Cys Val Ser Arg Asp Leu

660

665

670

Trp Cys Asp Gly Glu Ala Asp Cys Ser Asp Ser Ser Asp Glu Trp Asp

675

680

685

Cys Val Thr Leu Ser Ile Asn Val Asn Ser Ser Ser Phe Leu Met Val

690

695

700

His Arg Ala Ala Thr Glu His His Val Cys Ala Asp Gly Trp Gln Glu

705

710

715

720

Ile Leu Ser Gln Leu Ala Cys Lys Gln Met Gly Leu Gly Glu Pro Ser

725

730

735

Val Thr Lys Leu Ile Gln Glu Gln Glu Lys Glu Pro Arg Trp Leu Thr

740

745

750

Leu His Ser Asn Trp Glu Ser Leu Asn Gly Thr Thr Leu His Glu Leu

755

760

765

Leu Val Asn Gly Gln Ser Cys Glu Ser Arg Ser Lys Ile Ser Leu Leu

770

775

780

Cys Thr Lys Gln Asp Cys Gly Arg Arg Pro Ala Ala Arg Met Asn Lys  
 785 790 795 800

Arg Ile Leu Gly Gly Arg Thr Ser Arg Pro Gly Arg Trp Pro Trp Gln  
 805 810 815

Cys Ser Leu Gln Ser Glu Pro Ser Gly His Ile Cys Gly Cys Val Leu  
 820 825 830

Ile Ala Lys Lys Trp Val Leu Thr Val Ala His Cys Phe Glu Gly Arg  
 835 840 845

Glu Asn Ala Ala Val Trp Lys Val Val Leu Gly Ile Asn Asn Leu Asp  
 850 855 860

His Pro Ser Val Phe Met Gln Thr Arg Phe Val Lys Thr Ile Ile Leu  
 865 870 875 880

His Pro Arg Tyr Ser Arg Ala Val Val Asp Tyr Asp Ile Ser Ile Val  
 885 890 895

Glu Leu Ser Glu Asp Ile Ser Glu Thr Gly Tyr Val Arg Pro Val Cys  
 900 905 910

Leu Pro Asn Pro Glu Gln Trp Leu Glu Pro Asp Thr Tyr Cys Tyr Ile

915

920

925

Thr Gly Trp Gly His Met Gly Asn Lys Met Pro Phe Lys Leu Gln Glu

930

935

940

Gly Glu Val Arg Ile Ile Ser Leu Glu His Cys Gln Ser Tyr Phe Asp

945

950

955

960

Met Lys Thr Ile Thr Thr Arg Met Ile Cys Ala Gly Tyr Glu Ser Gly

965

970

975

Thr Val Asp Ser Cys Met Gly Asp Ser Gly Gly Pro Leu Val Cys Glu

980

985

990

Lys Pro Gly Gly Arg Trp Thr Leu Phe Gly Leu Thr Ser Trp Gly Ser

995

1000

1005

Val Cys Phe Ser Lys Val Leu Gly Pro Gly Val Tyr Ser Asn Val Ser

1010

1015

1020

Tyr Phe Val Glu Trp Ile Lys Arg Gln Ile Tyr Ile Gln Thr Phe Leu

1025

1030

1035

1040

Leu Asn

<210> 5

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<223> Description of Artificial Sequence:adapter for  
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26

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<213> Artificial Sequence

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cDNA amplification

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12

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26

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12



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26

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acacactcac ag

12

<210> 11

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ccagcatcga gaatcagtgt gacagt

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actgtcacac tg

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gtcgatgaac ttcgactgtc gatcgt

26

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acgatcgaca gt

12